



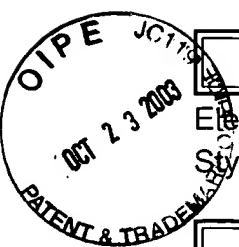
Electronic Filing System (EFS) Data
Electronic Patent Application Submission
USPTO Use Only

EFS ID: 49668
Application ID: 09743818
Title of Invention: Protease Susceptibility II
First Named Inventor: Anthony Weiss
Domestic/Foreign Application: Domestic Application
Filing Date: 2001-04-26
Effective Receipt Date: 2003-10-23
Submission Type: BIO Sequence Filing
Filing Type:
Confirmation number: 8602
Attorney Docket Number: GHC11USA




Total Fees Authorized:

Digital Certificate Holder: cn=Cathy A. Kodroff,ou=Registered Attorneys,ou=Patent and Trademark
Office,ou=Department of Commerce,o=U.S. Government,c=US
Certificate Message Digest: 79a77c537824f533a91104403ee220ce9152bff7



TRANSMITTAL

Electronic Version v1.1
Stylesheet Version v1.1.0

Title of Invention	Protease Susceptibility II											
Application Number: 09/743818 												
Date: 2001-04-26												
First Named Applicant: Anthony S. Weiss												
Confirmation Number: 8602												
Attorney Docket Number: GHC11USA												
<p>I hereby certify that the use of this system is for OFFICIAL correspondence between patent applicants or their representatives and the USPTO. Fraudulent or other use besides the filing of official correspondence by authorized parties is strictly prohibited, and subject to a fine and/or imprisonment under applicable law.</p> <p>I, the undersigned, certify that I have viewed a display of document(s) being electronically submitted to the United States Patent and Trademark Office, using either the USPTO provided style sheet or software, and that this is the document(s) I intend for initiation or further prosecution of a patent application noted in the submission. This document(s) will become part of the official electronic record at the USPTO.</p>												
<table border="1"><thead><tr><th>Submitted by:</th><th>Elec. Sign.</th><th>Sign. Capacity</th></tr></thead><tbody><tr><td>Cathy A. Kodroff Registered Number: 33,980</td><td>/cathyakodroff/</td><td>Attorney</td></tr></tbody></table>			Submitted by:	Elec. Sign.	Sign. Capacity	Cathy A. Kodroff Registered Number: 33,980	/cathyakodroff/	Attorney				
Submitted by:	Elec. Sign.	Sign. Capacity										
Cathy A. Kodroff Registered Number: 33,980	/cathyakodroff/	Attorney										
<table><tr><td>Documents being submitted</td><td>Files</td></tr><tr><td>us-bio-seq-trans</td><td>GHC11USA-usbios.xml</td></tr><tr><td></td><td>us-bio-seq-trans.dtd</td></tr><tr><td></td><td>us-bio-seq-trans.xsl</td></tr><tr><td>sequence-listing</td><td>sequence.txt</td></tr></table>			Documents being submitted	Files	us-bio-seq-trans	GHC11USA-usbios.xml		us-bio-seq-trans.dtd		us-bio-seq-trans.xsl	sequence-listing	sequence.txt
Documents being submitted	Files											
us-bio-seq-trans	GHC11USA-usbios.xml											
	us-bio-seq-trans.dtd											
	us-bio-seq-trans.xsl											
sequence-listing	sequence.txt											
Comments												



AMINO ACID AND/OR NUCLEOTIDE SEQUENCE LISTING SUBMISSION

Electronic Version v13

Stylesheet Version v01

This is a request for filing the electronic Computer Readable Form copy of a sequence listing via the Electronic Filing System for a patent application under 37 CFR 1.821-1.825 instead of via one of the physical media specified in 37 CFR 1.824(c).

This communication has an attached file which is an electronic copy of the amino acid and/or nucleotide sequence listing for the previously mentioned United States patent application.

The electronic copy submitted herewith is the Computer Readable Form (CRF), as required by 1.821(e).

Any applicable fees associated with the filing of the electronic copy have been paid.

This submission does not go beyond the disclosure of the application as originally filed (i.e., contains no new matter). It may be in addition to an original CRF, filed to comply with the sequence rules.

This submission in electronic form comprises only the CRF of 37 CFR 1.821(e). I acknowledge that I am responsible for all additional requirements of amino acid and/or nucleotide sequence listing submissions as specified in 37 CFR 1.821 - 1.825.

This submission does not go beyond the disclosure of the application as originally filed (i.e., contains no new matter), and/or is in addition to an original CRF filed to comply with the sequence rules. If not made to comply with an originally filed CRF, it is identical to the sequences disclosed in the application as originally filed and/or the paper copy of the sequence listing as originally filed.

I hereby certify that this correspondence is being transmitted to the United States Patent and Trademark Office on the following date: 2003-10-23

Name: Cathy A. Kodroff

Electronic Signature Mark: /CathyAKodroff/

Attachment
description:

Attached is a Substitute Sequence Listing. The hard copy and an appropriate extension of time is being supplied with a response to an Office Action dated 08/25/2003.

Compression

software used:



SEQUENCE LISTING

<110> Weiss, Anthony S.
<120> Protease Susceptibility II
<130> GHC11USA
<140> US 09/743,818
<141> 2001-04-26
<150> AU PP4723
<151> 1998-07-17
<150> PCT/AU99/00580
<151> 1999-07-19
<160> 105
<170> PatentIn version 3.2
<210> 1
<211> 2106
<212> DNA
<213> Homo sapiens

<400> 1
atgggtggcg ttccgggtgc tatcccggtt ggcgttccgg gtggtgtatt ctaccaggc 60
gcgggtctgg gtgcaactgg cggtgggtgc ctgggcccg gtggtaaacc gctgaaaccg 120
gttccaggcg gtctggcagg tgctggtctg ggtgcaggtc tgggcgcggt cccggcggtt 180
accttcccg gtgctctggt tccgggtggc gttgcagacg cagctgctgc gtacaaagcg 240
gcaaaggcag gtgcgggtct gggcggggta ccagggtgtg gcggtctggg tgtatctgct 300
ggcgcagttg ttccgcagcc ggggtgcagg gtaaaaccgg gcaaagttcc aggtgttggt 360
ctgccgggcg tatacccggtg tgggtgttctg ccgggcgcgc gtttccagg tggtggtgta 420
ctgccgggcg ttccgaccgg tgcagggtgt aaaccgaagg caccagggtg aggcggcgcg 480
ttcgcgggta tcccggtgtg tggcccggtt ggtggtccgc agccaggcgt tccgctgggt 540
taccgatca aagcgccgaa gcttccagg gtctacggtc tgccgtacac caccggtaaa 600
ctgccgtacg gctacggtcc ggggtggcgta gcagggtctg cgggtaaagc aggctaccca 660
accggtactg gtgttggtcc gcaggctgct gcggcagctg cggcgaaggc agcagcaaaa 720
ttcggcgcg gtgcagcggg tgttctgccg ggcgtagggt gtgctggcgt tccgggtgtt 780
ccagggtgca tcccgggcat cgggtgtatc gcaggcgtag gtactccggc ggccgctgcg 840
gctgcggcag ctgcggcgaa agcagctaaa tacggtgcgg cagcaggcct ggttccgggt 900
ggtccaggct tcgggtccgg tggtgtaggc gttccgggtg ctggtgttcc gggcgtaggt 960
gttccagggt cgggcattcc ggtgtgaccg ggtgcaggta tcccgggcgc tgcggttcca 1020

gggtgttgtat ccccggaagc ggcagctaag gctgctgcga aagctgcgaa atacggagct	1080
cgtccggggcg ttggtgttgg tggcatcccg acctacgggtg taggtgcagg cggtttccca	1140
ggtttcggcg ttggtgttgg tggcatcccg ggtgtagctg gtgttccgtc tgttggtggc	1200
gtaccgggtg ttggtggcgt tccaggtgta ggtatctccc cggaagcgca ggcagctgcg	1260
gcagctaaag cagcgaagta cggcgttggg actccggcgg cagcagctgc taaagcagcg	1320
gctaaagcag cgcagttcgg actagtcccg ggcgtaggtg ttgcgccagg tgttggcgta	1380
gcaccgggtg ttggtgttgc tccgggcgta ggtctggcac cgggtgttgg cgttgcacca	1440
gggtgtaggtg ttgcgccggg cgttgggtgta gcaccgggta tcgggtccggg tggcgttgcg	1500
gctgctgcga aatctgctgc gaaggttgct gcgaaagcgc agctgcgtgc agcagctggg	1560
ctgggtgcgg gcatcccagg tctgggtgta ggtgttgggtg ttccgggcct ggggtgtaggt	1620
gcaggggtac cgggcctggg tgttgggtgca ggcgttccgg gtttcgggtc tgttccgggc	1680
gcgctggctg ctgcgaaagc ggcgaaatac ggtgcagcgg ttccgggtgt actgggcggg	1740
ctgggtgctc tgggcgggtg ttgtatcccg ggcgggtgtg taggtgcagg cccagctgca	1800
gctgctgctg cggcaaaggc agcggcgaaa gcagctcagt tcgggtctggg tgggtgcagca	1860
gggtctgggcg gtctgggtgt tggcgggtctg ggtgtaccgg gcgttgggtg tctgggtggc	1920
atcccgccgg cggcggcagc taaagcggct aaatacgggtg cagcaggtct ggggtggcgtt	1980
ctgggtggtg ctgggtcagtt cccactgggc ggtgtagcgg cacgtccggg tttcgggtctg	2040
tccccgatct tcccaggcgg tgcattgcctg ggtaaagctt gcggccgtaa acgtaaataa	2100
tgatag	2106

<210> 2
 <211> 1992
 <212> DNA
 <213> Homo sapiens

<400> 2	
atgggtggcg ttccgggtgc tgttccgggt ggcgttccgg gtggtgtatt ctaccaggc	60
gcgggtttcg gtgctgttcc ggggtggcgtt gcagacgcag ctgctgcgta caaagcggca	120
aaggcagggtg cgggtctggg cggggtagca ggtgttggcg gtctgggtgt atctgctggc	180
gcagttgttc cgcagccggg tgcaggtgta aaaccgggca aagttccagg tgttgggtctg	240
ccgggcgtat acccgggttt cgggtgctgtt ccgggcgcgc gtttcccagg tgttgggtgta	300
ctgccgggcg ttccgaccgg tgcaggtggt aaaccgaagg caccaggtgt aggcggcgcg	360
ttcgcgggta tcccgggtgt tggcccgttc ggtggtccgc agccaggcgt tccgctgggt	420

tacccgatca aagcgccgaa gcttccaggt ggctacggtc tgccgtacac caccggtaaa	480
ctgccgtacg gctacgggtcc ggggtggcgta gcaggtgctg cgggtaaagc aggctacca	540
accggtactg gtgttgggtcc gcaggctgct gcggcagctg cggcgaaggc agcagcaaaa	600
ttcggcgcggt gtgcagcggg ttccgggtgct gttccggggcg taggtggtgc tggcgttccg	660
ggtgttccag gtgcgatccc gggcatcggg ggtatcgag gcgtaggtac tccggcggcc	720
gctgcggctg cggcagctgc ggcgaaagca gctaaatacg gtgcggcagc aggcctggtt	780
ccgggtggtc caggcttcgg tccgggtggt gtaggcgttc cgggtttcgg tgctgttccg	840
ggcgtaggtg ttccaggtgc gggcatcccg gttgtaccgg gtgcaggtat cccgggcgct	900
gcgggtttcg gtgctgtatc cccggaagcg gcagctaagg ctgctgcgaa agctgcgaaa	960
tacggagctc gtccgggctg tgggtgttgg ggcatcccga cctacggtgt aggtgcaggc	1020
ggtttccag gtttcggcgt tgggtgttgg ggcatcccgg gtgtagctgg tgttccgtct	1080
gttgggtggc taccgggtgt tgggtggcgt ccagggtgtag gtatctcccc ggaagcgag	1140
gcagctgcgg cagctaaagc agcgaagtac ggcgttggtg ctccggcggc agcagctgct	1200
aaagcagcgg ctaaagcagc gcagttcggg ctagttccgg gcgtaggtgt tgcgccaggt	1260
gttggcgtag caccgggtgt tgggtgttgc ccgggcgtag gtctggcacc ggggtgttggc	1320
gttgccagg gtgtaggtgt tgcgccgggc gttggtgtag caccgggtat cgggtccgggt	1380
ggcgttgcg ctgctgcgaa atctgctgcg aaggttgctg cgaaagcgca gctgcgtgca	1440
gcagctggtc tgggtgcggg catcccaggt ctgggtgtag gtgttggtgt tccgggcctg	1500
ggtgtaggtg caggggtacc gggcctgggt gttggtgcag gcgttccggg tttcgggtgct	1560
gttccggggc cgctggctgc tgcgaaagcg gcgaaatacg gtgctgttcc ggggtgtactg	1620
ggcgttctgg gtgctctggg cgggtgttgg atcccgggcg gtgttgtagg tgcaggccca	1680
gctgcagctg ctgctgcggc aaaggcagcg gcgaaagcag ctcagttcgg tctggttgggt	1740
gcagcaggtc tgggcgggtc ggggtgttgg ggtctgggtg taccgggcgt tgggtgtctg	1800
ggtggcatcc cgccggcggc ggcagctaaa gcggctaaat acggtgcagc aggtctgggt	1860
ggcgttctgg gtggtgctgg tcagttccca ctgggcgggt tagcggcacg tccgggtttc	1920
ggtctgtccc cgatcttccc aggcggtgca tgcctgggta aagcttgcg ccgtaaactg	1980
aaataatgat ag	1992

<210> 3
 <211> 2210
 <212> DNA
 <213> Homo sapiens

<220>
 <221> CDS
 <222> (3)..(2201)

<400> 3

ga tcc atg ggt ggc gtt ccg ggt gct atc ccg ggt ggc gtt ccg ggt	47
Ser Met Gly Gly Val Pro Gly Ala Ile Pro Gly Gly Val Pro Gly	
1 5 10 15	
ggt gta ttc tac cca ggc gcg ggt ctg ggt gca ctg ggc ggt ggt gcg	95
Gly Val Phe Tyr Pro Gly Ala Gly Leu Gly Ala Leu Gly Gly Gly Ala	
20 25 30	
ctg ggc ccg ggt ggt aaa ccg ctg aaa ccg gtt cca ggc ggt ctg gca	143
Leu Gly Pro Gly Gly Lys Pro Leu Lys Pro Val Pro Gly Gly Leu Ala	
35 40 45	
ggt gct ggt ctg ggt gca ggt ctg ggc gcg ttc ccg gcg gtt acc ttc	191
Gly Ala Gly Leu Gly Ala Gly Leu Gly Ala Phe Pro Ala Val Thr Phe	
50 55 60	
ccg ggt gct ctg gtt ccg ggt ggc gtt gca gac gca gct gct gcg tac	239
Pro Gly Ala Leu Val Pro Gly Gly Val Ala Asp Ala Ala Ala Tyr	
65 70 75	
aaa gcg gca aag gca ggt gcg ggt ctg ggc ggg gta cca ggt gtt ggc	287
Lys Ala Ala Lys Ala Gly Ala Gly Leu Gly Gly Val Pro Gly Val Gly	
80 85 90 95	
ggt ctg ggt gta tct gct ggc gca gtt gtt ccg cag ccg ggt gca ggt	335
Gly Leu Gly Val Ser Ala Gly Ala Val Val Pro Gln Pro Gly Ala Gly	
100 105 110	
gta aaa ccg ggc aaa gtt cca ggt gtt ggt ctg ccg ggc gta tac ccg	383
Val Lys Pro Gly Lys Val Pro Gly Val Gly Leu Pro Gly Val Tyr Pro	
115 120 125	
ggt ggt gtt ctg ccg ggc gcg cgt ttc cca ggt gtt ggt gta ctg ccg	431
Gly Gly Val Leu Pro Gly Ala Arg Phe Pro Gly Val Gly Val Leu Pro	
130 135 140	
ggc gtt ccg acc ggt gca ggt gtt aaa ccg aag gca cca ggt gta ggc	479
Gly Val Pro Thr Gly Ala Gly Val Lys Pro Lys Ala Pro Gly Val Gly	
145 150 155	
ggc gcg ttc gcg ggt atc ccg ggt gtt ggc ccg ttc ggt ggt ccg cag	527
Gly Ala Phe Ala Gly Ile Pro Gly Val Gly Pro Phe Gly Gly Pro Gln	
160 165 170 175	
cca ggc gtt ccg ctg ggt tac ccg atc aaa gcg ccg aag ctt cca ggt	575
Pro Gly Val Pro Leu Gly Tyr Pro Ile Lys Ala Pro Lys Leu Pro Gly	
180 185 190	
ggc tac ggt ctg ccg tac acc acc ggt aaa ctg ccg tac ggc tac ggt	623
Gly Tyr Gly Leu Pro Tyr Thr Thr Gly Lys Leu Pro Tyr Gly Tyr Gly	
195 200 205	
ccg ggt ggc gta gca ggt gct gcg ggt aaa gca ggc tac cca acc ggt	671
Pro Gly Gly Val Ala Gly Ala Ala Gly Lys Ala Gly Tyr Pro Thr Gly	

210	215	220	
act ggt gtt ggt ccg cag gct gct gcg gca gct gcg gcg aag gca gca Thr Gly Val Gly Pro Gln Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala 225 230 235			719
gca aaa ttc ggc gcg ggt gca gcg ggt gtt ctg ccg ggc gta ggt ggt Ala Lys Phe Gly Ala Gly Ala Ala Gly Val Leu Pro Gly Val Gly Gly 240 245 250 255			767
gct ggc gtt ccg ggt gtt cca ggt gcg atc ccg ggc atc ggt ggt atc Ala Gly Val Pro Gly Val Pro Gly Ala Ile Pro Gly Ile Gly Gly Ile 260 265 270			815
gca ggc gta ggt act ccg gcg gcc gct gcg gct gcg gca gct gcg gcg Ala Gly Val Gly Thr Pro Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala 275 280 285			863
aaa gca gct aaa tac ggt gcg gca gca ggc ctg gtt ccg ggt ggt cca Lys Ala Ala Lys Tyr Gly Ala Ala Ala Gly Leu Val Pro Gly Gly Pro 290 295 300			911
ggc ttc ggt ccg ggt gtt gta ggc gtt ccg ggt gct ggt gtt ccg ggc Gly Phe Gly Pro Gly Val Val Gly Val Pro Gly Ala Gly Val Pro Gly 305 310 315			959
gta ggt gtt cca ggt gcg ggc atc ccg gtt gta ccg ggt gca ggt atc Val Gly Val Pro Gly Ala Gly Ile Pro Val Val Pro Gly Ala Gly Ile 320 325 330 335			1007
ccg ggc gct gcg gtt cca ggt gtt gta tcc ccg gaa gcg gca gct aag Pro Gly Ala Ala Val Pro Gly Val Val Ser Pro Glu Ala Ala Ala Lys 340 345 350			1055
gct gct gcg aaa gct gcg aaa tac gga gct cgt ccg ggc gtt ggt gtt Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Arg Pro Gly Val Gly Val 355 360 365			1103
ggt ggc atc ccg acc tac ggt gta ggt gca ggc ggt ttc cca ggt ttc Gly Gly Ile Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe 370 375 380			1151
ggc gtt ggt gtt ggt ggc atc ccg ggt gta gct ggt gtt ccg tct gtt Gly Val Gly Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Ser Val 385 390 395			1199
ggt ggc gta ccg ggt gtt ggt ggc gtt cca ggt gta ggt atc tcc ccg Gly Gly Val Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro 400 405 410 415			1247
gaa gcg cag gca gct gcg gca gct aaa gca gcg aag tac ggc gtt ggt Glu Ala Gln Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly 420 425 430			1295
act ccg gcg gca gca gct gct aaa gca gcg gct aaa gca gcg cag ttc Thr Pro Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe 435 440 445			1343
gga cta gtt ccg ggc gta ggt gtt gcg cca ggt gtt ggc gta gca ccg Gly Leu Val Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro 450 455 460 465			1391

450	455	460	
ggt gtt ggt gtt gct ccg ggc gta ggt ctg gca ccg ggt gtt ggc gtt Gly Val Gly Val Ala Pro Gly Val Gly Leu Ala Pro Gly Val Gly Val 465 470 475			1439
gca cca ggt gta ggt gtt gcg ccg ggc gtt ggt gta gca ccg ggt atc Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Ile 480 485 490 495			1487
ggt ccg ggt ggc gtt gcg gct gct gcg aaa tct gct gcg aag gtt gct Gly Pro Gly Gly Val Ala Ala Ala Lys Ser Ala Ala Lys Val Ala 500 505 510			1535
gcg aaa gcg cag ctg cgt gca gca gct ggt ctg ggt gcg ggc atc cca Ala Lys Ala Gln Leu Arg Ala Ala Gly Leu Gly Ala Gly Ile Pro 515 520 525			1583
ggt ctg ggt gta ggt gtt ggt gtt ccg ggc ctg ggt gta ggt gca ggg Gly Leu Gly Val Gly Val Gly Val Pro Gly Leu Gly Val Gly Ala Gly 530 535 540			1631
gta ccg ggc ctg ggt gtt ggt gca ggc gtt ccg ggt ttc ggt gct ggc Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly Phe Gly Ala Gly 545 550 555			1679
gcg gac gaa ggt gta cgt cgt tcc ctg tct cca gaa ctg cgt gaa ggt Ala Asp Glu Gly Val Arg Arg Ser Leu Ser Pro Glu Leu Arg Glu Gly 560 565 570 575			1727
gac ccg tcc tct tcc cag cac ctg ccg tct acc ccg tcc tct cca cgt Asp Pro Ser Ser Ser Gln His Leu Pro Ser Thr Pro Ser Ser Pro Arg 580 585 590			1775
gtt ccg ggc gcg ctg gct gct gcg aaa gcg gcg aaa tac ggt gca gcg Val Pro Gly Ala Leu Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala 595 600 605			1823
gtt ccg ggt gta ctg ggc ggt ctg ggt gct ctg ggc ggt gtt ggt atc Val Pro Gly Val Leu Gly Gly Leu Gly Ala Leu Gly Gly Val Gly Ile 610 615 620			1871
ccg ggc ggt gtt gta ggt gca ggc cca gct gca gct gct gct gcg gca Pro Gly Gly Val Val Gly Ala Gly Pro Ala Ala Ala Ala Ala Ala 625 630 635			1919
aag gca gcg gcg aaa gca gct cag ttc ggt ctg gtt ggt gca gca ggt Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val Gly Ala Ala Gly 640 645 650 655			1967
ctg ggc ggt ctg ggt gtt ggc ggt ctg ggt gta ccg ggc gtt ggt ggt Leu Gly Gly Leu Gly Val Gly Gly Leu Gly Val Pro Gly Val Gly Gly 660 665 670			2015
ctg ggt ggc atc ccg ccg gcg gcg gca gct aaa gcg gct aaa tac ggt Leu Gly Gly Ile Pro Pro Ala Ala Ala Lys Ala Ala Lys Tyr Gly 675 680 685			2063
gca gca ggt ctg ggt ggc gtt ctg ggt ggt gct ggt cag ttc cca ctg Ala Ala Gly Leu Gly Gly Val Leu Gly Gly Ala Gly Gln Phe Pro Leu 2111			

690	695	700	
ggc ggt gta gcg gca cgt ccg ggt ttc ggt ctg tcc ccg atc ttc cca			2159
Gly Gly Val Ala Ala Arg Pro Gly Phe Gly Leu Ser Pro Ile Phe Pro			
705	710	715	
ggc ggt gca tgc ctg ggt aaa gct tgc ggc cgt aaa cgt aaa taatgatag			2210
Gly Gly Ala Cys Leu Gly Lys Ala Cys Gly Arg Lys Arg Lys			
720	725	730	
<210> 4			
<211> 733			
<212> PRT			
<213> Homo sapiens			
<400> 4			
Ser Met Gly Gly Val Pro Gly Ala Ile Pro Gly Gly Val Pro Gly Gly			
1	5	10	15
Val Phe Tyr Pro Gly Ala Gly Leu Gly Ala Leu Gly Gly Gly Ala Leu			
20	25	30	
Gly Pro Gly Gly Lys Pro Leu Lys Pro Val Pro Gly Gly Leu Ala Gly			
35	40	45	
Ala Gly Leu Gly Ala Gly Leu Gly Ala Phe Pro Ala Val Thr Phe Pro			
50	55	60	
Gly Ala Leu Val Pro Gly Gly Val Ala Asp Ala Ala Ala Ala Tyr Lys			
65	70	75	80
Ala Ala Lys Ala Gly Ala Gly Leu Gly Gly Val Pro Gly Val Gly Gly			
85	90	95	
Leu Gly Val Ser Ala Gly Ala Val Val Pro Gln Pro Gly Ala Gly Val			
100	105	110	
Lys Pro Gly Lys Val Pro Gly Val Gly Leu Pro Gly Val Tyr Pro Gly			
115	120	125	
Gly Val Leu Pro Gly Ala Arg Phe Pro Gly Val Gly Val Leu Pro Gly			
130	135	140	
Val Pro Thr Gly Ala Gly Val Lys Pro Lys Ala Pro Gly Val Gly Gly			
145	150	155	160
Ala Phe Ala Gly Ile Pro Gly Val Gly Pro Phe Gly Gly Pro Gln Pro			
165	170	175	

Gly Val Pro Leu Gly Tyr Pro Ile Lys Ala Pro Lys Leu Pro Gly Gly
 180 185 190

Tyr Gly Leu Pro Tyr Thr Thr Gly Lys Leu Pro Tyr Gly Tyr Gly Pro
 195 200 205

Gly Gly Val Ala Gly Ala Ala Gly Lys Ala Gly Tyr Pro Thr Gly Thr
 210 215 220

Gly Val Gly Pro Gln Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala
 225 230 235 240

Lys Phe Gly Ala Gly Ala Ala Gly Val Leu Pro Gly Val Gly Gly Ala
 245 250 255

Gly Val Pro Gly Val Pro Gly Ala Ile Pro Gly Ile Gly Gly Ile Ala
 260 265 270

Gly Val Gly Thr Pro Ala Ala Ala Ala Ala Ala Ala Ala Ala Lys
 275 280 285

Ala Ala Lys Tyr Gly Ala Ala Ala Gly Leu Val Pro Gly Gly Pro Gly
 290 295 300

Phe Gly Pro Gly Val Val Gly Val Pro Gly Ala Gly Val Pro Gly Val
 305 310 315 320

Gly Val Pro Gly Ala Gly Ile Pro Val Val Pro Gly Ala Gly Ile Pro
 325 330 335

Gly Ala Ala Val Pro Gly Val Val Ser Pro Glu Ala Ala Ala Lys Ala
 340 345 350

Ala Ala Lys Ala Ala Lys Tyr Gly Ala Arg Pro Gly Val Gly Val Gly
 355 360 365

Gly Ile Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe Gly
 370 375 380

Val Gly Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Ser Val Gly
 385 390 395 400

Gly Val Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro Glu
 405 410 415

Ala Gln Ala Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly Thr
 420 425 430

Pro Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly
 435 440 445

Leu Val Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly
 450 455 460

Val Gly Val Ala Pro Gly Val Gly Leu Ala Pro Gly Val Gly Val Ala
 465 470 475 480

Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Ile Gly
 485 490 495

Pro Gly Gly Val Ala Ala Ala Ala Lys Ser Ala Ala Lys Val Ala Ala
 500 505 510

Lys Ala Gln Leu Arg Ala Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly
 515 520 525

Leu Gly Val Gly Val Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val
 530 535 540

Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly Phe Gly Ala Gly Ala
 545 550 555 560

Asp Glu Gly Val Arg Arg Ser Leu Ser Pro Glu Leu Arg Glu Gly Asp
 565 570 575

Pro Ser Ser Ser Gln His Leu Pro Ser Thr Pro Ser Ser Pro Arg Val
 580 585 590

Pro Gly Ala Leu Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Val
 595 600 605

Pro Gly Val Leu Gly Gly Leu Gly Ala Leu Gly Gly Val Gly Ile Pro
 610 615 620

Gly Gly Val Val Gly Ala Gly Pro Ala Ala Ala Ala Ala Ala Ala Lys
 625 630 635 640

Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val Gly Ala Ala Gly Leu
 645 650 655

Gly Gly Leu Gly Val Gly Gly Leu Gly Val Pro Gly Val Gly Gly Leu
660 665 670

Gly Gly Ile Pro Pro Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala
675 680 685

Ala Gly Leu Gly Gly Val Leu Gly Gly Ala Gly Gln Phe Pro Leu Gly
690 695 700

Gly Val Ala Ala Arg Pro Gly Phe Gly Leu Ser Pro Ile Phe Pro Gly
705 710 715 720

Gly Ala Cys Leu Gly Lys Ala Cys Gly Arg Lys Arg Lys
725 730

<210> 5
<211> 698
<212> PRT
<213> Homo sapiens

<400> 5

Gly Gly Val Pro Gly Ala Ile Pro Gly Gly Val Pro Gly Gly Val Phe
1 5 10 15

Tyr Pro Gly Ala Gly Leu Gly Ala Leu Gly Gly Gly Ala Leu Gly Pro
20 25 30

Gly Gly Lys Pro Leu Lys Pro Val Pro Gly Gly Leu Ala Gly Ala Gly
35 40 45

Leu Gly Ala Gly Leu Gly Ala Phe Pro Ala Val Thr Phe Pro Gly Ala
50 55 60

Leu Val Pro Gly Gly Val Ala Asp Ala Ala Ala Tyr Lys Ala Ala
65 70 75 80

Lys Ala Gly Ala Gly Leu Gly Gly Val Pro Gly Val Gly Gly Leu Gly
85 90 95

Val Ser Ala Gly Ala Val Val Pro Gln Pro Gly Ala Gly Val Lys Pro
100 105 110

Gly Lys Val Pro Gly Val Gly Leu Pro Gly Val Tyr Pro Gly Gly Val
115 120 125

Leu Pro Gly Ala Arg Phe Pro Gly Val Gly Val Leu Pro Gly Val Pro
 130 135 140

Thr Gly Ala Gly Val Lys Pro Lys Ala Pro Gly Val Gly Gly Ala Phe
 145 150 155 160

Ala Gly Ile Pro Gly Val Gly Pro Phe Gly Gly Pro Gln Pro Gly Val
 165 170 175

Pro Leu Gly Tyr Pro Ile Lys Ala Pro Lys Leu Pro Gly Gly Tyr Gly
 180 185 190

Leu Pro Tyr Thr Thr Gly Lys Leu Pro Tyr Gly Tyr Gly Pro Gly Gly
 195 200 205

Val Ala Gly Ala Ala Gly Lys Ala Gly Tyr Pro Thr Gly Thr Gly Val
 210 215 220

Gly Pro Gln Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Phe
 225 230 235 240

Gly Ala Gly Ala Ala Gly Val Leu Pro Gly Val Gly Gly Ala Gly Val
 245 250 255

Pro Gly Val Pro Gly Ala Ile Pro Gly Ile Gly Gly Ile Ala Gly Val
 260 265 270

Gly Thr Pro Ala Ala Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala
 275 280 285

Lys Tyr Gly Ala Ala Ala Gly Leu Val Pro Gly Gly Pro Gly Phe Gly
 290 295 300

Pro Gly Val Val Gly Val Pro Gly Ala Gly Val Pro Gly Val Gly Val
 305 310 315 320

Pro Gly Ala Gly Ile Pro Val Val Pro Gly Ala Gly Ile Pro Gly Ala
 325 330 335

Ala Val Pro Gly Val Val Ser Pro Glu Ala Ala Ala Lys Ala Ala Ala
 340 345 350

Lys Ala Ala Lys Tyr Gly Ala Arg Pro Gly Val Gly Val Gly Gly Ile
 355 360 365

Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe Gly Val Gly
 370 375 380

Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Ser Val Gly Gly Val
 385 390 395 400

Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro Glu Ala Gln
 405 410 415

Ala Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly Thr Pro Ala
 420 425 430

Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val
 435 440 445

Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly
 450 455 460

Val Ala Pro Gly Val Gly Leu Ala Pro Gly Val Gly Val Ala Pro Gly
 465 470 475 480

Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Ile Gly Pro Gly
 485 490 495

Gly Val Ala Ala Ala Ala Lys Ser Ala Ala Lys Val Ala Ala Lys Ala
 500 505 510

Gln Leu Arg Ala Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly Leu Gly
 515 520 525

Val Gly Val Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly
 530 535 540

Leu Gly Val Gly Ala Gly Val Pro Gly Phe Gly Ala Val Pro Gly Ala
 545 550 555 560

Leu Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Val Pro Gly Val
 565 570 575

Leu Gly Gly Leu Gly Ala Leu Gly Gly Val Gly Ile Pro Gly Gly Val
 580 585 590

Val Gly Ala Gly Pro Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala
 595 600 605

Lys Ala Ala Gln Phe Gly Leu Val Gly Ala Ala Gly Leu Gly Gly Leu
610 615 620

Gly Val Gly Gly Leu Gly Val Pro Gly Val Gly Gly Leu Gly Gly Ile
625 630 635 640

Pro Pro Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Gly Leu
645 650 655

Gly Gly Val Leu Gly Gly Ala Gly Gln Phe Pro Leu Gly Gly Val Ala
660 665 670

Ala Arg Pro Gly Phe Gly Leu Ser Pro Ile Phe Pro Gly Gly Ala Cys
675 680 685

Leu Gly Lys Ala Cys Gly Arg Lys Arg Lys
690 695

<210> 6
<211> 661
<212> PRT
<213> Homo sapiens

<400> 6

Met Gly Gly Val Pro Gly Ala Val Pro Gly Gly Val Pro Gly Gly Val
1 5 10 15

Phe Tyr Pro Gly Ala Gly Phe Gly Ala Val Pro Gly Gly Val Ala Asp
20 25 30

Ala Ala Ala Ala Tyr Lys Ala Ala Lys Ala Gly Ala Gly Leu Gly Gly
35 40 45

Val Pro Gly Val Gly Gly Leu Gly Val Ser Ala Gly Ala Val Val Pro
50 55 60

Gln Pro Gly Ala Gly Val Lys Pro Gly Lys Val Pro Gly Val Gly Leu
65 70 75 80

Pro Gly Val Tyr Pro Gly Phe Gly Ala Val Pro Gly Ala Arg Phe Pro
85 90 95

Gly Val Gly Val Leu Pro Gly Val Pro Thr Gly Ala Gly Val Lys Pro
100 105 110

Lys Ala Pro Gly Val Gly Gly Ala Phe Ala Gly Ile Pro Gly Val Gly
115 120 125

Pro Phe Gly Gly Pro Gln Pro Gly Val Pro Leu Gly Tyr Pro Ile Lys
130 135 140

Ala Pro Lys Leu Pro Gly Gly Tyr Gly Leu Pro Tyr Thr Thr Gly Lys
145 150 155 160

Leu Pro Tyr Gly Tyr Gly Pro Gly Gly Val Ala Gly Ala Ala Gly Lys
165 170 175

Ala Gly Tyr Pro Thr Gly Thr Gly Val Gly Pro Gln Ala Ala Ala Ala
180 185 190

Ala Ala Ala Lys Ala Ala Ala Lys Phe Gly Ala Gly Ala Ala Gly Phe
195 200 205

Gly Ala Val Pro Gly Val Gly Gly Ala Gly Val Pro Gly Val Pro Gly
210 215 220

Ala Ile Pro Gly Ile Gly Gly Ile Ala Gly Val Gly Thr Pro Ala Ala
225 230 235 240

Ala Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala
245 250 255

Ala Gly Leu Val Pro Gly Gly Pro Gly Phe Gly Pro Gly Val Val Gly
260 265 270

Val Pro Gly Phe Gly Ala Val Pro Gly Val Gly Val Pro Gly Ala Gly
275 280 285

Ile Pro Val Val Pro Gly Ala Gly Ile Pro Gly Ala Ala Gly Phe Gly
290 295 300

Ala Val Ser Pro Glu Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Lys
305 310 315 320

Tyr Gly Ala Arg Pro Gly Val Gly Val Gly Gly Ile Pro Thr Tyr Gly
325 330 335

Val Gly Ala Gly Gly Phe Pro Gly Phe Gly Val Gly Val Gly Gly Ile
340 345 350

Pro Gly Val Ala Gly Val Pro Ser Val Gly Gly Val Pro Gly Val Gly
 355 360 365

Gly Val Pro Gly Val Gly Ile Ser Pro Glu Ala Gln Ala Ala Ala Ala
 370 375 380

Ala Lys Ala Ala Lys Tyr Gly Val Gly Thr Pro Ala Ala Ala Ala Ala
 385 390 395 400

Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val Pro Gly Val Gly
 405 410 415

Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly
 420 425 430

Val Gly Leu Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala
 435 440 445

Pro Gly Val Gly Val Ala Pro Gly Ile Gly Pro Gly Gly Val Ala Ala
 450 455 460

Ala Ala Lys Ser Ala Ala Lys Val Ala Ala Lys Ala Gln Leu Arg Ala
 465 470 475 480

Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly Leu Gly Val Gly Val Gly
 485 490 495

Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly Leu Gly Val Gly
 500 505 510

Ala Gly Val Pro Gly Phe Gly Ala Val Pro Gly Ala Leu Ala Ala Ala
 515 520 525

Lys Ala Ala Lys Tyr Gly Ala Val Pro Gly Val Leu Gly Gly Leu Gly
 530 535 540

Ala Leu Gly Gly Val Gly Ile Pro Gly Gly Val Val Gly Ala Gly Pro
 545 550 555 560

Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe
 565 570 575

Gly Leu Val Gly Ala Ala Gly Leu Gly Gly Leu Gly Val Gly Gly Leu
 580 585 590

Gly Val Pro Gly Val Gly Gly Leu Gly Gly Ile Pro Pro Ala Ala Ala
595 600 605

Ala Lys Ala Ala Lys Tyr Gly Ala Ala Gly Leu Gly Gly Val Leu Gly
610 615 620

Gly Ala Gly Gln Phe Pro Leu Gly Gly Val Ala Ala Arg Pro Gly Phe
625 630 635 640

Gly Leu Ser Pro Ile Phe Pro Gly Gly Ala Cys Leu Gly Lys Ala Cys
645 650 655

Gly Arg Lys Arg Lys
660

<210> 7
<211> 571
<212> PRT
<213> Homo sapiens

<400> 7

Gly Gly Val Pro Gly Ala Ile Pro Gly Gly Val Pro Gly Gly Val Phe
1 5 10 15

Tyr Pro Gly Ala Gly Leu Gly Ala Leu Gly Gly Gly Ala Leu Gly Pro
20 25 30

Gly Gly Lys Pro Leu Lys Pro Val Pro Gly Gly Leu Ala Gly Ala Gly
35 40 45

Leu Gly Ala Gly Leu Gly Ala Phe Pro Ala Val Thr Phe Pro Gly Ala
50 55 60

Leu Val Pro Gly Gly Val Ala Asp Ala Ala Ala Tyr Lys Ala Ala
65 70 75 80

Lys Ala Gly Ala Gly Leu Gly Gly Val Pro Gly Val Gly Gly Leu Gly
85 90 95

Val Ser Ala Gly Ala Val Val Pro Gln Pro Gly Ala Gly Val Lys Pro
100 105 110

Gly Lys Val Pro Gly Val Gly Leu Pro Gly Val Tyr Pro Gly Gly Val
115 120 125

Leu Pro Gly Ala Arg Phe Pro Gly Val Gly Val Leu Pro Gly Val Pro

130		135		140
Thr Gly Ala Gly Val Lys Pro Lys Ala Pro Gly Val Gly Gly Ala Phe				
145		150		155 160
Ala Gly Ile Pro Gly Val Gly Pro Phe Gly Gly Pro Gln Pro Gly Val				
	165		170	175
Pro Leu Gly Tyr Pro Ile Lys Ala Pro Lys Leu Pro Gly Gly Tyr Gly				
	180		185	190
Leu Pro Tyr Thr Thr Gly Lys Leu Pro Tyr Gly Tyr Gly Pro Gly Gly				
	195		200	205
Val Ala Gly Ala Ala Gly Lys Ala Gly Tyr Pro Thr Gly Thr Gly Val				
	210		215	220
Gly Pro Gln Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Phe				
225		230		235 240
Gly Ala Gly Ala Ala Gly Val Leu Pro Gly Val Gly Gly Ala Gly Val				
	245		250	255
Pro Gly Val Pro Gly Ala Ile Pro Gly Ile Gly Gly Ile Ala Gly Val				
	260		265	270
Gly Thr Pro Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala				
	275		280	285
Lys Tyr Gly Ala Ala Ala Gly Leu Val Pro Gly Gly Pro Gly Phe Gly				
	290		295	300
Pro Gly Val Val Gly Val Pro Gly Ala Gly Val Pro Gly Val Gly Val				
305		310		315 320
Pro Gly Ala Gly Ile Pro Val Val Pro Gly Ala Gly Ile Pro Gly Ala				
	325		330	335
Ala Val Pro Gly Val Val Ser Pro Glu Ala Ala Ala Lys Ala Ala Ala				
	340		345	350
Lys Ala Ala Lys Tyr Gly Ala Arg Pro Gly Val Gly Val Gly Gly Ile				
	355		360	365
Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe Gly Val Gly				

370 375 380
 Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Ser Val Gly Gly Val
 385 390 395 400
 Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro Glu Ala Gln
 405 410 415
 Ala Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly Thr Pro Ala
 420 425 430
 Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val
 435 440 445
 Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly
 450 455 460
 Val Ala Pro Gly Val Gly Leu Ala Pro Gly Val Gly Val Ala Pro Gly
 465 470 475 480
 Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Ile Gly Pro Gly
 485 490 495
 Gly Val Ala Ala Ala Ala Lys Ser Ala Ala Lys Val Ala Ala Lys Ala
 500 505 510
 Gln Leu Arg Ala Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly Leu Gly
 515 520 525
 Val Gly Val Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly
 530 535 540
 Leu Gly Val Gly Ala Gly Cys Ser Gly Phe Arg Cys Trp Arg Gly Arg
 545 550 555 560
 Arg Cys Thr Ser Phe Pro Val Ser Arg Thr Ala
 565 570

<210> 8
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 8

Lys Ala Pro Gly Val Gly Gly Ala Phe
 1 5

<210> 9
<211> 7
<212> PRT
<213> Homo sapiens

<400> 9

Arg Ala Ala Ala Gly Leu Gly
1 5

<210> 10
<211> 11
<212> PRT
<213> Homo sapiens

<400> 10

Arg Ser Leu Ser Pro Glu Leu Arg Glu Gly Asp
1 5 10

<210> 11
<211> 9
<212> PRT
<213> Homo sapiens

<400> 11

Lys Ala Ala Lys Ala Gly Ala Gly Leu
1 5

<210> 12
<211> 9
<212> PRT
<213> Homo sapiens

<400> 12

Lys Ala Gly Ala Gly Leu Gly Gly Val
1 5

<210> 13
<211> 13
<212> PRT
<213> Homo sapiens

<400> 13

Ala Leu Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala
1 5 10

<210> 14
<211> 11
<212> PRT

<213> Homo sapiens

<400> 14

Lys Ala Ala Gln Phe Gly Leu Val Pro Gly Val
1 5 10

<210> 15

<211> 11

<212> PRT

<213> Homo sapiens

<400> 15

Lys Ser Ala Ala Lys Val Ala Ala Lys Ala Gln
1 5 10

<210> 16

<211> 9

<212> PRT

<213> Homo sapiens

<400> 16

Arg Ser Leu Ser Pro Glu Leu Arg Glu
1 5

<210> 17

<211> 8

<212> PRT

<213> Homo sapiens

<400> 17

Gly Gln Leu Arg Ala Ala Ala Gly
1 5

<210> 18

<211> 8

<212> PRT

<213> Homo sapiens

<400> 18

Val Gln Leu Arg Ala Ala Ala Gly
1 5

<210> 19

<211> 8

<212> PRT

<213> Homo sapiens

<400> 19

Ile Gln Leu Arg Ala Ala Ala Gly

1 5

<210> 20
<211> 8
<212> PRT
<213> Homo sapiens

<400> 20

Leu Gln Leu Arg Ala Ala Ala Gly
1 5

<210> 21
<211> 8
<212> PRT
<213> Homo sapiens

<400> 21

Ala Asn Leu Arg Ala Ala Ala Gly
1 5

<210> 22
<211> 8
<212> PRT
<213> Homo sapiens

<400> 22

Ala Gly Leu Arg Ala Ala Ala Gly
1 5

<210> 23
<211> 8
<212> PRT
<213> Homo sapiens

<400> 23

Ala Val Leu Arg Ala Ala Ala Gly
1 5

<210> 24
<211> 8
<212> PRT
<213> Homo sapiens

<400> 24

Ala Ser Leu Arg Ala Ala Ala Gly
1 5

<210> 25
<211> 8

<212> PRT
<213> Homo sapiens

<400> 25

Ala Gln Gly Arg Ala Ala Ala Gly
1 5

<210> 26
<211> 8
<212> PRT
<213> Homo sapiens

<400> 26

Ala Gln Val Arg Ala Ala Ala Gly
1 5

<210> 27
<211> 8
<212> PRT
<213> Homo sapiens

<400> 27

Ala Gln Ile Arg Ala Ala Ala Gly
1 5

<210> 28
<211> 8
<212> PRT
<213> Homo sapiens

<400> 28

Ala Gln Ala Arg Ala Ala Ala Gly
1 5

<210> 29
<211> 8
<212> PRT
<213> Homo sapiens

<400> 29

Ala Gln Leu Arg Gly Ala Ala Gly
1 5

<210> 30
<211> 8
<212> PRT
<213> Homo sapiens

<400> 30

Ala Gln Leu Arg Val Ala Ala Gly
1 5

<210> 31
<211> 8
<212> PRT
<213> Homo sapiens

<400> 31

Ala Gln Leu Arg Ile Ala Ala Gly
1 5

<210> 32
<211> 8
<212> PRT
<213> Homo sapiens

<400> 32

Ala Gln Leu Arg Leu Ala Ala Gly
1 5

<210> 33
<211> 8
<212> PRT
<213> Homo sapiens

<400> 33

Ala Gln Leu Arg Ala Gly Ala Gly
1 5

<210> 34
<211> 8
<212> PRT
<213> Homo sapiens

<400> 34

Ala Gln Leu Arg Ala Val Ala Gly
1 5

<210> 35
<211> 8
<212> PRT
<213> Homo sapiens

<400> 35

Ala Gln Leu Arg Ala Ile Ala Gly
1 5

<210> 36

<211> 8
<212> PRT
<213> Homo sapiens

<400> 36

Ala Gln Leu Arg Ala Leu Ala Gly
1 5

<210> 37
<211> 8
<212> PRT
<213> Homo sapiens

<400> 37

Ala Gln Leu Arg Ala Ala Gly Gly
1 5

<210> 38
<211> 8
<212> PRT
<213> Homo sapiens

<400> 38

Ala Gln Leu Arg Ala Ala Val Gly
1 5

<210> 39
<211> 8
<212> PRT
<213> Homo sapiens

<400> 39

Ala Gln Leu Arg Ala Ala Ile Gly
1 5

<210> 40
<211> 8
<212> PRT
<213> Homo sapiens

<400> 40

Ala Gln Leu Arg Ala Ala Leu Gly
1 5

<210> 41
<211> 8
<212> PRT
<213> Homo sapiens

<400> 41

Ala Gln Leu Arg Ala Ala Ala Ala
1 5

<210> 42
<211> 8
<212> PRT
<213> Homo sapiens

<400> 42

Ala Gln Leu Arg Ala Ala Ala Ile
1 5

<210> 43
<211> 8
<212> PRT
<213> Homo sapiens

<400> 43

Ala Gln Leu Arg Ala Ala Ala Val
1 5

<210> 44
<211> 8
<212> PRT
<213> Homo sapiens

<400> 44

Ala Gln Leu Arg Ala Ala Ala Leu
1 5

<210> 45
<211> 8
<212> PRT
<213> Homo sapiens

<400> 45

Val Gly Gly Ala Leu Ala Ala Ala
1 5

<210> 46
<211> 8
<212> PRT
<213> Homo sapiens

<400> 46

Gly Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 47
<211> 8
<212> PRT
<213> Homo sapiens

<400> 47

Ile Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 48
<211> 8
<212> PRT
<213> Homo sapiens

<400> 48

Leu Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 49
<211> 8
<212> PRT
<213> Homo sapiens

<400> 49

Ala Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 50
<211> 8
<212> PRT
<213> Homo sapiens

<400> 50

Val Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 51
<211> 8
<212> PRT
<213> Homo sapiens

<400> 51

Val Pro Ile Ala Leu Ala Ala Ala
1 5

<210> 52
<211> 8
<212> PRT
<213> Homo sapiens

<400> 52

Val Pro Leu Ala Leu Ala Ala Ala
1 5

<210> 53

<211> 8

<212> PRT

<213> Homo sapiens

<400> 53

Val Pro Val Ala Leu Ala Ala Ala
1 5

<210> 54

<211> 8

<212> PRT

<213> Homo sapiens

<400> 54

Val Pro Gly Ala Gly Ala Ala Ala
1 5

<210> 55

<211> 8

<212> PRT

<213> Homo sapiens

<400> 55

Val Pro Gly Ala Ile Ala Ala Ala
1 5

<210> 56

<211> 8

<212> PRT

<213> Homo sapiens

<400> 56

Val Pro Gly Ala Ala Ala Ala Ala
1 5

<210> 57

<211> 8

<212> PRT

<213> Homo sapiens

<400> 57

Val Pro Gly Ala Val Ala Ala Ala
1 5

<210> 58
<211> 8
<212> PRT
<213> Homo sapiens

<400> 58

Val Pro Gly Ala Leu Gly Ala Ala
1 5

<210> 59
<211> 8
<212> PRT
<213> Homo sapiens

<400> 59

Val Pro Gly Ala Leu Ile Ala Ala
1 5

<210> 60
<211> 8
<212> PRT
<213> Homo sapiens

<400> 60

Val Pro Gly Ala Leu Leu Ala Ala
1 5

<210> 61
<211> 8
<212> PRT
<213> Homo sapiens

<400> 61

Val Pro Gly Ala Leu Val Ala Ala
1 5

<210> 62
<211> 8
<212> PRT
<213> Homo sapiens

<400> 62

Val Pro Gly Ala Leu Ala Gly Ala
1 5

<210> 63
<211> 8
<212> PRT
<213> Homo sapiens

<400> 63

Val Pro Gly Ala Leu Ala Ile Ala
1 5

<210> 64

<211> 8

<212> PRT

<213> Homo sapiens

<400> 64

Val Pro Gly Ala Leu Ala Leu Ala
1 5

<210> 65

<211> 8

<212> PRT

<213> Homo sapiens

<400> 65

Val Pro Gly Ala Leu Ala Val Ala
1 5

<210> 66

<211> 8

<212> PRT

<213> Homo sapiens

<400> 66

Val Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 67

<211> 8

<212> PRT

<213> Homo sapiens

<400> 67

Val Pro Gly Ala Leu Ala Ala Gly
1 5

<210> 68

<211> 8

<212> PRT

<213> Homo sapiens

<400> 68

Val Pro Gly Ala Leu Ala Ala Ile
1 5

<210> 69
<211> 8
<212> PRT
<213> Homo sapiens

<400> 69

Val Pro Gly Ala Leu Ala Ala Leu
1 5

<210> 70
<211> 8
<212> PRT
<213> Homo sapiens

<400> 70

Val Pro Gly Ala Leu Ala Ala Val
1 5

<210> 71
<211> 515
<212> PRT
<213> Homo sapiens

<400> 71

Gly Gly Val Pro Gly Ala Ile Pro Gly Gly Val Pro Gly Gly Val Phe
1 5 10 15

Tyr Pro Gly Ala Gly Leu Gly Ala Leu Gly Gly Gly Ala Leu Gly Pro
20 25 30

Gly Gly Lys Pro Leu Lys Pro Val Pro Gly Gly Leu Ala Gly Ala Gly
35 40 45

Leu Gly Ala Gly Leu Gly Ala Phe Pro Ala Val Thr Phe Pro Gly Ala
50 55 60

Leu Val Pro Gly Gly Val Ala Asp Ala Ala Ala Tyr Lys Ala Ala
65 70 75 80

Lys Ala Gly Ala Gly Leu Gly Gly Val Pro Gly Val Gly Gly Leu Gly
85 90 95

Val Ser Ala Gly Ala Val Val Pro Gln Pro Gly Ala Gly Val Lys Pro
100 105 110

Gly Lys Val Pro Gly Val Gly Leu Pro Gly Val Tyr Pro Gly Gly Val

115		120		125
Leu Pro Gly Ala Arg Phe	Pro Gly Val Gly Val	Leu Pro Gly Val Pro		
130	135	140		
Thr Gly Ala Gly Val Lys	Pro Lys Ala Pro Gly Val	Gly Gly Ala Phe		
145	150	155	160	
Ala Gly Ile Pro Gly Val	Gly Pro Phe Gly Gly Pro Gln	Pro Gly Val		
	165	170	175	
Pro Leu Gly Tyr Pro Ile	Lys Ala Pro Lys Leu Pro	Gly Gly Tyr Gly		
	180	185	190	
Leu Pro Tyr Thr Thr Gly	Lys Leu Pro Tyr Gly Tyr	Gly Pro Gly Gly		
	195	200	205	
Val Ala Gly Ala Ala Gly	Lys Ala Gly Tyr Pro Thr	Gly Thr Gly Val		
	210	215	220	
Gly Pro Gln Ala Ala Ala	Ala Ala Ala Ala Lys Ala	Ala Ala Lys Phe		
225	230	235	240	
Gly Ala Gly Ala Ala Gly	Val Leu Pro Gly Val Gly	Gly Ala Gly Val		
	245	250	255	
Pro Gly Val Pro Gly Ala	Ile Pro Gly Ile Gly Gly	Ile Ala Gly Val		
	260	265	270	
Gly Thr Pro Ala Ala Ala	Ala Ala Ala Ala Ala	Ala Lys Ala Ala		
	275	280	285	
Lys Tyr Gly Ala Ala Ala	Gly Leu Val Pro Gly Gly	Pro Gly Phe Gly		
	290	295	300	
Pro Gly Val Val Gly Val	Pro Gly Ala Gly Val	Pro Gly Val Gly Val		
305	310	315	320	
Pro Gly Ala Gly Ile	Pro Val Val Pro Gly Ala	Gly Ile Pro Gly Ala		
	325	330	335	
Ala Val Pro Gly Val Val	Ser Pro Glu Ala Ala	Lys Ala Ala Ala		
	340	345	350	
Lys Ala Ala Lys Tyr Gly	Ala Arg Pro Gly Val Gly	Val Gly Gly Ile		

355 360 365
 Pro Thr Tyr Gly Val Gly Ala Gly Gly Phe Pro Gly Phe Gly Val Gly
 370 375 380
 Val Gly Gly Ile Pro Gly Val Ala Gly Val Pro Ser Val Gly Gly Val
 385 390 395 400
 Pro Gly Val Gly Gly Val Pro Gly Val Gly Ile Ser Pro Glu Ala Gln
 405 410 415
 Ala Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Val Gly Thr Pro Ala
 420 425 430
 Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala Gln Phe Gly Leu Val
 435 440 445
 Pro Gly Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Val Gly
 450 455 460
 Val Ala Pro Gly Val Gly Leu Ala Pro Gly Val Gly Val Ala Pro Gly
 465 470 475 480
 Val Gly Val Ala Pro Gly Val Gly Val Ala Pro Gly Ile Gly Pro Gly
 485 490 495
 Gly Val Ala Ala Ala Ala Lys Ser Ala Ala Lys Val Ala Ala Lys Ala
 500 505 510
 Gln Leu Arg
 515

<210> 72
 <211> 49
 <212> PRT
 <213> Homo sapiens

<400> 72

Ala Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly Leu Gly Val Gly Val
 1 5 10 15
 Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly Leu Gly Val
 20 25 30
 Gly Ala Gly Val Pro Gly Phe Gly Ala Gly Ala Asp Glu Gly Val Arg
 35 40 45

Arg

<210> 73
<211> 171
<212> PRT
<213> Homo sapiens

<400> 73

Gly Val Arg Arg Ser Leu Ser Pro Glu Leu Arg Glu Gly Asp Pro Ser
1 5 10 15

Ser Ser Gln His Leu Pro Ser Thr Pro Ser Ser Pro Arg Val Pro Gly
20 25 30

Ala Leu Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Val Pro Gly
35 40 45

Val Leu Gly Gly Leu Gly Ala Leu Gly Gly Val Gly Ile Pro Gly Gly
50 55 60

Val Val Gly Ala Gly Pro Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala
65 70 75 80

Ala Lys Ala Ala Gln Phe Gly Leu Val Gly Ala Ala Gly Leu Gly Gly
85 90 95

Leu Gly Val Gly Gly Leu Gly Val Pro Gly Val Gly Gly Leu Gly Gly
100 105 110

Ile Pro Pro Ala Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Gly
115 120 125

Leu Gly Gly Val Leu Gly Gly Ala Gly Gln Phe Pro Leu Gly Gly Val
130 135 140

Ala Ala Arg Pro Gly Phe Gly Leu Ser Pro Ile Phe Pro Gly Gly Ala
145 150 155 160

Cys Leu Gly Lys Ala Cys Gly Arg Lys Arg Lys
165 170

<210> 74
<211> 183
<212> PRT

<213> Homo sapiens

<400> 74

Ala Ala Ala Gly Leu Gly Ala Gly Ile Pro Gly Leu Gly Val Gly Val
1 5 10 15

Gly Val Pro Gly Leu Gly Val Gly Ala Gly Val Pro Gly Leu Gly Val
20 25 30

Gly Ala Gly Val Pro Gly Phe Gly Ala Val Pro Gly Ala Leu Ala Ala
35 40 45

Ala Lys Ala Ala Lys Tyr Gly Ala Ala Val Pro Gly Val Leu Gly Gly
50 55 60

Leu Gly Ala Leu Gly Gly Val Gly Ile Pro Gly Gly Val Val Gly Ala
65 70 75 80

Gly Pro Ala Ala Ala Ala Ala Ala Ala Lys Ala Ala Ala Lys Ala Ala
85 90 95

Gln Phe Gly Leu Val Gly Ala Ala Gly Leu Gly Gly Leu Gly Val Gly
100 105 110

Gly Leu Gly Val Pro Gly Val Gly Gly Leu Gly Gly Ile Pro Pro Ala
115 120 125

Ala Ala Ala Lys Ala Ala Lys Tyr Gly Ala Ala Gly Leu Gly Gly Val
130 135 140

Leu Gly Gly Ala Gly Gln Phe Pro Leu Gly Gly Val Ala Ala Arg Pro
145 150 155 160

Gly Phe Gly Leu Ser Pro Ile Phe Pro Gly Gly Ala Cys Leu Gly Lys
165 170 175

Ala Cys Gly Arg Lys Arg Lys
180

<210> 75

<211> 18

<212> PRT

<213> bovine tropoelastin

<400> 75

Val Pro Thr Gly Ala Gly Val Lys Pro Lys Ala Pro Gly Gly Gly Gly

1 5 10 15

Ala Phe

<210> 76
<211> 17
<212> PRT
<213> mouse tropoelastin

<400> 76

Val Pro Thr Gly Thr Gly Val Lys Ala Lys Ala Pro Gly Gly Gly Ala
1 5 10 15

Phe

<210> 77
<211> 18
<212> PRT
<213> bovine elastin

<400> 77

Val Pro Thr Gly Ala Gly Val Lys Pro Lys Ala Gln Val Gly Ala Gly
1 5 10 15

Ala Phe

<210> 78
<211> 16
<212> PRT
<213> rat tropoelastin

<400> 78

Val Pro Thr Gly Thr Gly Val Lys Ala Lys Val Pro Gly Gly Gly Gly
1 5 10 15

<210> 79
<211> 15
<212> PRT
<213> chicken tropoelastin

<400> 79

Val Pro Thr Gly Thr Gly Ile Lys Ala Lys Gly Pro Gly Ala Gly
1 5 10 15

<210> 80

<211> 17
<212> PRT
<213> mouse tropoelastin

<400> 80

Lys Ala Ala Ala Lys Ala Gln Tyr Arg Ala Ala Ala Gly Leu Gly Ala
1 5 10 15

Gly

<210> 81
<211> 17
<212> PRT
<213> bovine elastin

<400> 81

Lys Ala Ala Ala Lys Ala Gln Phe Arg Ala Ala Ala Gly Leu Pro Ala
1 5 10 15

Gly

<210> 82
<211> 20
<212> PRT
<213> Artificial

<220>
<223> tropoelastin consensus sequence

<220>
<221> MISC_FEATURE
<222> (9)..(9)
<223> IS AN AROMATIC OR HYDROPHOBIC RESIDUE

<220>
<221> MISC_FEATURE
<222> (16)..(16)
<223> can be either Pro or Gly

<220>
<221> MISC_FEATURE
<222> (19)..(19)
<223> is a hydrophobic residue

<400> 82

Ala Lys Ala Ala Ala Lys Ala Gln Xaa Arg Ala Ala Ala Gly Leu Xaa
1 5 10 15

Ala Gly Xaa Pro

<210> 83
 <211> 14
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> (7)..(8)
 <223> there is a reduced peptide bond between Arg and Ala

<400> 83

Ala Ala Lys Ala Gln Leu Arg Ala Ala Ala Gly Leu Gly Ala
 1 5 10

<210> 84
 <211> 14
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> (7)..(8)
 <223> there is a reduced peptide bond between Ala and Arg

<400> 84

Ala Gly Leu Gly Ala Ala Ala Arg Leu Gln Ala Lys Ala Ala
 1 5 10

<210> 85
 <211> 14
 <212> PRT
 <213> Homo sapiens

<400> 85

Ala Gly Leu Gly Ala Ala Ala Arg Leu Gln Ala Lys Ala Ala
 1 5 10

<210> 86
 <211> 8
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> (4)..(5)
 <223> there is a reduced peptide bond between Ala and Leu

<400> 86

Val Pro Gly Ala Leu Ala Ala Ala
1 5

<210> 87
<211> 8
<212> PRT
<213> Homo sapiens

<220>
<221> VARIANT
<222> (4)..(5)
<223> there is a reduced peptide bond between Leu and Ala

<400> 87

Ala Ala Ala Leu Ala Gly Pro Val
1 5

<210> 88
<211> 8
<212> PRT
<213> Homo sapiens

<400> 88

Ala Ala Ala Leu Ala Gly Pro Val
1 5

<210> 89
<211> 30
<212> DNA
<213> Artificial

<220>
<223> mutagenic primer

<400> 89
cggggtttcgg tgctgttccg ggcgcgctgg

30

<210> 90
<211> 20
<212> DNA
<213> Artificial

<220>
<223> primer

<400> 90
gggtggtggc gttgcaccag

20

<210> 91
<211> 20
<212> DNA

<213> Artificial
 <220>
 <223> primer
 <400> 91
 tgcacctaca acaccgcccg 20
 <210> 92
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 92
 tgcctttgcc ggtttgtacg 20
 <210> 93
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 93
 tccaggtggc tacggtctgc 20
 <210> 94
 <211> 21
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 94
 gagtacctac gcctgcgata c 21
 <210> 95
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 95
 ggagtaccaa cgccgtactt 20
 <210> 96
 <211> 20
 <212> DNA

<213> Artificial
 <220>
 <223> primer
 <400> 96
 ggggtgttggc gttgcaccag 20
 <210> 97
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 97
 tgcacctaca acaccgccccg 20
 <210> 98
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 98
 gcactcacta tagggagacc 20
 <210> 99
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 99
 gccaaactcag cttcctttcg 20
 <210> 100
 <211> 20
 <212> DNA
 <213> Artificial
 <220>
 <223> primer
 <400> 100
 taatacgact cactataggg 20
 <210> 101
 <211> 15
 <212> PRT

<213> Homo sapiens

<400> 101

Val	Val	Gly	Ser	Pro	Ser	Ala	Gln	Asp	Glu	Ala	Ser	Pro	Leu	Ser
1				5					10					15

<210> 102

<211> 10

<212> PRT

<213> Homo sapiens

<400> 102

Lys	Ala	Ala	Ala	Lys	Ala	Gly	Ala	Gly	Leu
1				5					10

<210> 103

<211> 12

<212> PRT

<213> Homo sapiens

<400> 103

Ala	Leu	Ala	Ala	Lys	Ala	Ala	Lys	Tyr	Gly	Ala	Ala
1				5					10		

<210> 104

<211> 11

<212> PRT

<213> Homo sapiens

<400> 104

Lys	Ala	Ala	Gln	Phe	Gly	Leu	Val	Pro	Gly	Val
1				5					10	

<210> 105

<211> 18

<212> PRT

<213> Homo sapiens

<400> 105

Gly	Gly	Val	Pro	Gly	Ala	Ile	Pro	Gly	Gly	Val	Pro	Gly	Gly	Phe	Tyr
1				5					10					15	

Pro Gly